

SEQUENCE LISTING

<110> INSTITUT PASTEUR

<120> PKNB KINASE AND PSTP PHOSPHATASE AND METHODS OF IDENTIFYING
INHIBITORY SUBSTANCES

<130> D22486

<150> US 60/487,943

<151> 2003-07-18

<160> 20

<170> PatentIn version 3.2

<210> 1

<211> 514

<212> PRT

<213> Mycobacterium tuberculosis

<400> 1

Met	Ala	Arg	Val	Thr	Leu	Val	Leu	Arg	Tyr	Ala	Ala	Arg	Ser	Asp	Arg
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Gly	Leu	Val	Arg	Ala	Asn	Asn	Glu	Asp	Ser	Val	Tyr	Ala	Gly	Ala	Arg
		20						25					30		

Leu	Leu	Ala	Leu	Ala	Asp	Gly	Met	Gly	Gly	His	Ala	Ala	Gly	Glu	Val
		35					40					45			

Ala	Ser	Gln	Leu	Val	Ile	Ala	Ala	Leu	Ala	His	Leu	Asp	Asp	Asp	Glu
	50					55					60				

Pro	Gly	Gly	Asp	Leu	Leu	Ala	Lys	Leu	Asp	Ala	Ala	Val	Arg	Ala	Gly
65					70					75					80

Asn	Ser	Ala	Ile	Ala	Ala	Gln	Val	Glu	Met	Glu	Pro	Asp	Leu	Glu	Gly
			85						90					95	

Met	Gly	Thr	Thr	Leu	Thr	Ala	Ile	Leu	Phe	Ala	Gly	Asn	Arg	Leu	Gly
			100					105					110		

Leu	Val	His	Ile	Gly	Asp	Ser	Arg	Gly	Tyr	Leu	Leu	Arg	Asp	Gly	Glu
		115					120					125			

Leu	Thr	Gln	Ile	Thr	Lys	Asp	Asp	Thr	Phe	Val	Gln	Thr	Leu	Val	Asp
	130					135					140				

Glu	Gly	Arg	Ile	Thr	Pro	Glu	Glu	Ala	His	Ser	His	Pro	Gln	Arg	Ser
145					150					155					160

Leu	Ile	Met	Arg	Ala	Leu	Thr	Gly	His	Glu	Val	Glu	Pro	Thr	Leu	Thr
				165					170					175	

Met	Arg	Glu	Ala	Arg	Ala	Gly	Asp	Arg	Tyr	Leu	Leu	Cys	Ser	Asp	Gly
			180					185					190		

Leu	Ser	Asp	Pro	Val	Ser	Asp	Glu	Thr	Ile	Leu	Glu	Ala	Leu	Gln	Ile
		195					200					205			

Pro Glu Val Ala Glu Ser Ala His Arg Leu Ile Glu Leu Ala Leu Arg
 210 215 220
 Gly Gly Gly Pro Asp Asn Val Thr Val Val Val Ala Asp Val Val Asp
 225 230 235 240
 Tyr Asp Tyr Gly Gln Thr Gln Pro Ile Leu Ala Gly Ala Val Ser Gly
 245 250 255
 Asp Asp Asp Gln Leu Thr Leu Pro Asn Thr Ala Ala Gly Arg Ala Ser
 260 265 270
 Ala Ile Ser Gln Arg Lys Glu Ile Val Lys Arg Val Pro Pro Gln Ala
 275 280 285
 Asp Thr Phe Ser Arg Pro Arg Trp Ser Gly Arg Arg Leu Ala Phe Val
 290 295 300
 Val Ala Leu Val Thr Val Leu Met Thr Ala Gly Leu Leu Ile Gly Arg
 305 310 315 320
 Ala Ile Ile Arg Ser Asn Tyr Tyr Val Ala Asp Tyr Ala Gly Ser Val
 325 330 335
 Ser Ile Met Arg Gly Ile Gln Gly Ser Leu Leu Gly Met Ser Leu His
 340 345 350
 Gln Pro Tyr Leu Met Gly Cys Leu Ser Pro Arg Asn Glu Leu Ser Gln
 355 360 365
 Ile Ser Tyr Gly Gln Ser Gly Gly Pro Leu Asp Cys His Leu Met Lys
 370 375 380
 Leu Glu Asp Leu Arg Pro Pro Glu Arg Ala Gln Val Arg Ala Gly Leu
 385 390 395 400
 Pro Ala Gly Thr Leu Asp Asp Ala Ile Gly Gln Leu Arg Glu Leu Ala
 405 410 415
 Ala Asn Ser Leu Leu Pro Pro Cys Pro Ala Pro Arg Ala Thr Ser Pro
 420 425 430
 Pro Gly Arg Pro Ala Pro Pro Thr Thr Ser Glu Thr Thr Glu Pro Asn
 435 440 445
 Val Thr Ser Ser Pro Ala Ser Pro Ser Pro Thr Thr Ser Ala Pro Ala
 450 455 460
 Pro Thr Gly Thr Thr Pro Ala Ile Pro Thr Ser Ala Ser Pro Ala Ala
 465 470 475 480
 Pro Ala Ser Pro Pro Thr Pro Trp Pro Val Thr Ser Ser Pro Thr Met
 485 490 495
 Ala Ala Leu Pro Pro Pro Pro Pro Gln Pro Gly Ile Asp Cys Arg Ala
 500 505 510
 Ala Ala

<210> 2
 <211> 382
 <212> PRT
 <213> Homo sapiens

<400> 2

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Met Gly Ala Phe Leu Asp Lys Pro Lys Met Glu Lys His Asn Ala Gln
1           5           10           15

Gly Gln Gly Asn Gly Leu Arg Tyr Gly Leu Ser Ser Met Gln Gly Trp
          20           25           30

Arg Val Glu Met Glu Asp Ala His Thr Ala Val Ile Gly Leu Pro Ser
          35           40           45

Gly Leu Glu Ser Trp Ser Phe Phe Ala Val Tyr Asp Gly His Ala Gly
          50           55           60

Ser Gln Val Ala Lys Tyr Cys Cys Glu His Leu Leu Asp His Ile Thr
65           70           75           80

Asn Asn Gln Asp Phe Lys Gly Ser Ala Gly Ala Pro Ser Val Glu Asn
          85           90           95

Val Lys Asn Gly Ile Arg Thr Gly Phe Leu Glu Ile Asp Glu His Met
          100          105          110

Arg Val Met Ser Glu Lys Lys His Gly Ala Asp Arg Ser Gly Ser Thr
          115          120          125

Ala Val Gly Val Leu Ile Ser Pro Gln His Thr Tyr Phe Ile Asn Cys
          130          135          140

Gly Asp Ser Arg Gly Leu Leu Cys Arg Asn Arg Lys Val His Phe Phe
145          150          155          160

Thr Gln Asp His Lys Pro Ser Asn Pro Leu Glu Lys Glu Arg Ile Gln
          165          170          175

Asn Ala Gly Gly Ser Val Met Ile Gln Arg Val Asn Gly Ser Leu Ala
          180          185          190

Val Ser Arg Ala Leu Gly Asp Phe Asp Tyr Lys Cys Val His Gly Lys
          195          200          205

Gly Pro Thr Glu Gln Leu Val Ser Pro Glu Pro Glu Val His Asp Ile
          210          215          220

Glu Arg Ser Glu Glu Asp Asp Gln Phe Ile Ile Leu Ala Cys Asp Gly
225          230          235          240

Ile Trp Asp Val Met Gly Asn Glu Glu Leu Cys Asp Phe Val Arg Ser
          245          250          255

Arg Leu Glu Val Thr Asp Asp Leu Glu Lys Val Cys Asn Glu Val Val
          260          265          270

Asp Thr Cys Leu Tyr Lys Gly Ser Arg Asp Asn Met Ser Val Ile Leu
          275          280          285

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Ile Cys Phe Pro Asn Ala Pro Lys Val Ser Pro Glu Ala Val Lys Lys
 290 295 300

Glu Ala Glu Leu Asp Lys Tyr Leu Glu Cys Arg Val Glu Glu Ile Ile
 305 310 315 320

Lys Lys Gln Gly Glu Gly Val Pro Asp Leu Val His Val Met Arg Thr
 325 330 335

Leu Ala Ser Glu Asn Ile Pro Ser Leu Pro Pro Gly Gly Glu Leu Ala
 340 345 350

Ser Lys Arg Asn Val Ile Glu Ala Val Tyr Asn Arg Leu Asn Pro Tyr
 355 360 365

Lys Asn Asp Asp Thr Asp Ser Thr Ser Thr Asp Asp Met Trp
 370 375 380

<210> 3

<211> 271

<212> PRT

<213> Mycobacterium tuberculosis

<400> 3

Ile Thr Arg Asp Val Gln Val Pro Asp Val Arg Gly Gln Ser Ser Ala
 1 5 10 15

Asp Ala Ile Ala Thr Leu Gln Asn Arg Gly Phe Lys Ile Arg Thr Leu
 20 25 30

Gln Lys Pro Asp Ser Thr Ile Pro Pro Asp His Val Ile Gly Thr Asp
 35 40 45

Pro Ala Ala Asn Thr Ser Val Ser Ala Gly Asp Glu Ile Thr Val Asn
 50 55 60

Val Ser Thr Gly Pro Glu Gln Arg Glu Ile Pro Asp Val Ser Thr Leu
 65 70 75 80

Thr Tyr Ala Glu Ala Val Lys Lys Leu Thr Ala Ala Gly Phe Gly Arg
 85 90 95

Phe Lys Gln Ala Asn Ser Pro Ser Thr Pro Glu Leu Val Gly Lys Val
 100 105 110

Ile Gly Thr Asn Pro Pro Ala Asn Gln Thr Ser Ala Ile Thr Asn Val
 115 120 125

Val Ile Ile Ile Val Gly Ser Gly Pro Ala Thr Lys Asp Ile Pro Asp
 130 135 140

Val Ala Gly Gln Thr Val Asp Val Ala Gln Lys Asn Leu Asn Val Tyr
 145 150 155 160

Gly Phe Thr Lys Phe Ser Gln Ala Ser Val Asp Ser Pro Arg Pro Ala
 165 170 175

Gly Glu Val Thr Gly Thr Asn Pro Pro Ala Gly Thr Thr Val Pro Val

Pro Asp Leu Ser Gly Met Phe Trp Val Asp Ala Glu Pro Arg Leu Arg
210 215 220

Ala Leu Gly Trp Thr Gly Met Leu Asp Lys Gly Ala Asp Val Asp Ala
225 230 235 240

Gly Gly Ser Gln His Asn Arg Val Val Tyr Gln Asn Pro Pro Ala Gly
245 250 255

Thr Gly Val Asn Arg Asp Gly Ile Ile Thr Leu Arg Phe Gly Gln
260 265 270

<210> 4

<211> 271

<212> PRT

<213> Mycobacterium leprae

<400> 4

Asn Thr Arg Asp Val Gln Val Pro Asp Val Arg Gly Gln Val Ser Ala
1 5 10 15

Asp Ala Ile Ser Ala Leu Gln Asn Arg Gly Phe Lys Thr Arg Thr Leu
20 25 30

Gln Lys Pro Asp Ser Thr Ile Pro Pro Asp His Val Ile Ser Thr Glu
35 40 45

Pro Gly Ala Asn Ala Ser Val Gly Ala Gly Asp Glu Ile Thr Ile Asn
50 55 60

Val Ser Thr Gly Pro Glu Gln Arg Glu Val Pro Asp Val Ser Ser Leu
65 70 75 80

Asn Tyr Thr Asp Ala Val Lys Lys Leu Thr Ser Ser Gly Phe Lys Ser
85 90 95

Phe Lys Gln Ala Asn Ser Pro Ser Thr Pro Glu Leu Leu Gly Lys Val
100 105 110

```
Ile Gly Thr Asn Pro Ser Ala Asn Gln Thr Ser Ala Ile Thr Asn Val  
          115                      120                      125
```

Ile Thr Ile Ile Val Gly Ser Gly Pro Glu Thr Lys Gln Ile Pro Asp
130 135 140

Val	Thr	Gly	Gln	Ile	Val	Glu	Ile	Ala	Gln	Lys	Asn	Leu	Asn	Val	Tyr
145					150					155					160

Gly Phe Thr Lys Phe Ser Gln Ala Ser Val Asp Ser Pro Arg Pro Thr
165 170 175

Gly Glu Val Ile Gly Thr Asn Pro Pro Lys Asp Ala Thr Val Pro Val

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180							185					190			
Asp	Ser	Val	Ile	Glu	Leu	Gln	Val	Ser	Lys	Gly	Asn	Gln	Phe	Val	Met
		195					200					205			
Pro	Asp	Leu	Ser	Gly	Met	Phe	Trp	Ala	Asp	Ala	Glu	Pro	Arg	Leu	Arg
	210					215					220				
Ala	Leu	Gly	Trp	Thr	Gly	Val	Leu	Asp	Lys	Gly	Pro	Asp	Val	Asp	Ala
225					230					235					240
Gly	Gly	Ser	Gln	His	Asn	Arg	Val	Ala	Tyr	Gln	Asn	Pro	Pro	Ala	Gly
				245					250					255	
Ala	Gly	Val	Asn	Arg	Asp	Gly	Ile	Ile	Thr	Leu	Lys	Phe	Gly	Gln	
			260				265					270			

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<210> 5
<211> 274
<212> PRT
<213> Corynebacterium glutamicum
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<400> 5

Ser	Thr	Ala	Thr	Ser	Ala	Ile	Pro	Asn	Val	Glu	Gly	Leu	Pro	Gln	Gln
1				5				10						15	
Glu	Ala	Leu	Thr	Glu	Leu	Gln	Ala	Ala	Gly	Phe	Val	Val	Asn	Ile	Val
			20					25					30		
Glu	Glu	Ala	Ser	Ala	Asp	Val	Ala	Glu	Gly	Leu	Val	Ile	Arg	Ala	Asn
		35					40					45			
Pro	Ser	Val	Gly	Ser	Glu	Ile	Arg	Gln	Gly	Ala	Thr	Val	Thr	Ile	Thr
	50					55					60				
Val	Ser	Thr	Gly	Arg	Glu	Met	Ile	Asn	Ile	Pro	Asp	Val	Ser	Gly	Met
65					70					75					80
Thr	Leu	Glu	Asp	Ala	Ala	Arg	Ala	Leu	Glu	Asp	Val	Gly	Leu	Ile	Leu
				85					90					95	
Asn	Gln	Asn	Val	Arg	Glu	Glu	Thr	Ser	Asp	Asp	Val	Glu	Ser	Gly	Leu
			100					105					110		
Val	Ile	Asp	Gln	Asn	Pro	Glu	Ala	Gly	Gln	Glu	Val	Val	Val	Gly	Ser
		115					120					125			
Ser	Val	Ser	Leu	Thr	Met	Ser	Ser	Gly	Thr	Glu	Ser	Ile	Arg	Val	Pro
	130					135					140				
Asn	Leu	Thr	Gly	Met	Asn	Trp	Ser	Gln	Ala	Glu	Gln	Asn	Leu	Ile	Ser
145					150					155					160
Met	Gly	Phe	Asn	Pro	Thr	Ala	Ser	Tyr	Leu	Asp	Ser	Ser	Glu	Pro	Glu
				165					170					175	
Gly	Glu	Val	Leu	Ser	Val	Ser	Ser	Gln	Gly	Thr	Glu	Leu	Pro	Lys	Gly
			180					185					190		

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Ser Ser Ile Thr Val Glu Val Ser Asn Gly Met Leu Ile Gln Ala Pro
 195 200 205

Asp Leu Ala Arg Met Ser Thr Glu Gln Ala Ile Ser Ala Leu Arg Ala
 210 215 220

Ala Gly Trp Thr Ala Pro Asp Gln Ser Leu Ile Val Gly Asp Pro Ile
 225 230 235 240

His Thr Ala Ala Leu Val Asp Gln Asn Lys Ile Gly Phe Gln Ser Pro
 245 250 255

Thr Pro Ala Thr Leu Phe Arg Lys Asp Ala Gln Val Gln Val Arg Leu
 260 265 270

Phe Glu

<210> 6

<211> 268

<212> PRT

<213> Thermobifida fusca

<400> 6

Gly Arg Tyr Glu Thr Val Pro Asp Leu Val Gly Val Glu Ser Asp Glu
 1 5 10 15

Ala Arg Arg Asp Leu Arg Met Leu Gly Phe Arg Val Gln Thr Ala Glu
 20 25 30

Glu Pro Ala Tyr Ser Asp Glu Ala Pro Pro Gly Thr Val Ala Ala Thr
 35 40 45

Asp Pro Glu Ala Gly Ser Arg Leu Leu Pro Asp Thr Leu Val Thr Leu
 50 55 60

Ile Leu Ser Ala Gly Pro Gln Tyr Val Glu Met Pro Asp Val Glu Gly
 65 70 75 80

Ala Ser Val Ala Glu Ala Arg Asp Ala Leu Lys Glu Val Gly Leu Thr
 85 90 95

Asp Ile Val Glu Asp Glu Ile Thr Ser Phe Asp Asn Pro Pro Gly Thr
 100 105 110

Val Ile Thr Thr Lys Pro Ala Pro Gly Glu Lys Ala Asn Arg Glu Glu
 115 120 125

Ser Val Thr Leu Thr Ile Ser Ala Gly Phe Pro Met Pro Asn Val Val
 130 135 140

Gly Gln Lys Val Asp Asp Ala Arg Arg Leu Leu Glu Ser Ser Asp Leu
 145 150 155 160

Glu Val Thr Val Val Glu Glu His His Asp Glu Val Pro Glu Gly His
 165 170 175

Val Ile Ser Gln Glu Pro Glu Ala Glu Thr Thr Val Gly Ala Gly Gln
 180 185 190

Ser Val Thr Leu Thr Val Ser Ser Gly Pro Glu Leu Val Glu Val Pro
 195 200 205

Asp Ile Arg Gly Trp Lys Val Asp Lys Ala Arg Lys Glu Leu Glu Glu
 210 215 220

Arg Gly Phe Glu Val Thr Val His Gln Val Ile Gly Asn Arg Val Gly
 225 230 235 240

Asp Tyr Asn Pro Lys Gly Glu Ala Pro Lys Gly Ser Thr Ile Glu Ile
 245 250 255

Trp Thr Ser Pro Phe Gly Arg Glu Arg Asp Arg Asp
 260 265

<210> 7

<211> 274

<212> PRT

<213> Corynebacterium efficiens

<400> 7

Ser Ala Ser Thr Gln Gln Ile Pro Asn Ile Val Gly Leu Pro Glu Asn
 1 5 10 15

Glu Ala Val Leu Glu Leu Glu Arg Leu Gly Phe Thr Val Val Leu Thr
 20 25 30

Thr Glu Pro Ser Pro Asp Val Ala Glu Gly Leu Val Ile Arg Thr Ser
 35 40 45

Pro Asn Val Gly Ser Glu Ile Arg Glu Gly Ala Thr Val Thr Leu Thr
 50 55 60

Ile Ser Ser Gly Arg Glu Val Val Thr Ile Pro Asp Val Thr Gly Leu
 65 70 75 80

Thr Leu Ala Glu Ala Thr Arg Glu Ile Glu Gly Ala Gly Leu Val Leu
 85 90 95

Asp Gln Ser Ile Arg Glu Glu Asn Ser Asp Asp Tyr Pro Ala Gly Thr
 100 105 110

Val Ile Gln Gln Asn Pro Arg Ala Gly Gly Glu Thr Ser Val Gly Ala
 115 120 125

Ser Ile Thr Leu Thr Val Ser Thr Gly Pro Ser Leu Val Arg Val Pro
 130 135 140

Val Ile Thr Gly Met Gln Trp Ser Gln Ala Glu Ser Asn Ile Thr Ser
 145 150 155 160

Leu Gly Leu Val Pro Asp Ile Tyr Tyr Val Asp Ser Leu Leu Pro Glu
 165 170 175

Gly Gln Val Ile Ser Ala Ser Gly Gln Gly Thr Glu Leu Pro Arg Gly
 180 185 190

Ser Thr Val Thr Val Glu Ile Ser Asn Gly Met Leu Ile Glu Ala Pro

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195	200	205
Asp Leu Ala Arg Leu Asp Val Asp Asn Ala Leu Lys Ala Leu Arg Asp		
210	215	220
Ala Gly Trp Thr Ala Pro Asp Thr Ser Leu Ile Glu Gly Ala Pro Ile		
225	230	235 240
Pro Thr Gly Ala Leu Val Asp Gln Gly Arg Ile Gly Phe Gln Asp Pro		
	245	250 255
Ser Pro Gly Gln Pro Leu Arg Lys Asp Ala Val Val Asn Ile Arg Leu		
	260	265 270
Tyr Arg		

<210> 8
 <211> 212
 <212> PRT
 <213> Thermobifida fusca

<400> 8

Gly Thr Asp Asn Ile Thr Ile Pro Asn Val Ala Gly Met Ser Val Glu		
1	5	10 15
Glu Ala Thr Glu Thr Leu Gln Glu Lys Gly Phe Glu Asn Ile Glu Val		
	20	25 30
Ala Asp Glu Pro Thr Pro Ser Asn Glu Ile Glu Glu Gly Lys Val Val		
	35	40 45
Gly Thr Asp Pro Glu Ile Gly Glu Thr Val Pro Pro Asp Thr Glu Ile		
	50	55 60
Thr Ile Leu Ile Ser Gly Gly Pro Glu Met Ile Glu Met Pro Asp Leu		
	65	70 75 80
Val Gly Met Ser Gln Ala Asp Ala Leu Gly Glu Ile Asn Arg Ala Gly		
	85	90 95
Leu Ala Arg Gly Glu Ile Thr His Gln Glu Ser Asp Glu Pro Gln Gly		
	100	105 110
Thr Val Leu Ser Thr Asp Pro Lys Ala Gly Thr Glu Val Glu Pro Gly		
	115	120 125
Thr Lys Val Asn Leu Val Val Ala Lys Ala Ser Thr Lys Val Glu Val		
	130	135 140
Pro Ser Leu Ala Gly Met Asn Glu Asp Gln Ala Arg Glu Arg Leu Ala		
	145	150 155 160
Glu Leu Gly Leu Thr Leu Glu Ala Gln Thr Gln Glu Thr Ser Asp Ala		
	165	170 175
Thr Pro Gly Thr Ala Ile Ala Gln Ser Pro Gln Ala Gly Thr Lys Val		
	180	185 190

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Glu Arg Gly Thr Thr Val Thr Val Thr Phe Ala Lys Glu Pro Gln Arg
 195 200 205

Pro Glu Pro Pro
 210

<210> 9

<211> 278

<212> PRT

<213> Bifidobacterium longum

<400> 9

Ser Glu Asp Thr Val Thr Ile Pro Glu Val Cys Asn Ala Ser Thr Ser
 1 5 10 15

Lys Asp Ser Ile Glu Leu Lys Leu Lys Ala Ser Gly Leu Lys Met Thr
 20 25 30

Glu Lys Gln Asp Thr Asp Ser Thr Glu Pro Glu Gly Thr Cys Thr Lys
 35 40 45

Met Ser Pro Asp Ala Gly Ser Lys Val Ala Lys Gly Ser Ala Val Lys
 50 55 60

Val Trp Phe Ser Ala Gly Pro Gln Ser Thr Gln Val Pro Asp Val Lys
 65 70 75 80

Glu Arg Ser Gln Glu Glu Ala Arg Ser Ile Leu Glu Ser Ala Gly Phe
 85 90 95

Lys Val Asn Ala Ala Val Lys Thr Glu Asp Ser Ala Asp Ile Ala Lys
 100 105 110

Asp Met Val Thr Lys Thr Asp Pro Ala Ala Gly Gln Ser Val Pro Lys
 115 120 125

Gly Thr Thr Ile Thr Ile Tyr Val Ser Ser Gly Met Thr Thr Val Pro
 130 135 140

Ser Asn Leu Val Gly Gln Ser Lys Asp Ser Val Leu Gln Gln Tyr Glu
 145 150 155 160

Gly Lys Phe Ser Phe Thr Val Glu Gln Glu Ser Ser Asp Thr Val Glu
 165 170 175

Ala Gly Leu Ile Thr Arg Val Ser Pro Asp Ser Gly Ser Ser Ile Ala
 180 185 190

Gln Gly Gly Phe Ile Thr Ile Trp Val Ser Thr Gly Lys Glu Lys Val
 195 200 205

Ala Val Pro Asn Ile Thr Ala Gly Thr Asp Tyr Val Thr Ala Glu Leu
 210 215 220

Met Leu Lys Ala Val Gly Leu Lys Ala Gln Ala Asn Gly Pro Thr Gly
 225 230 235 240

Ser Thr Ala Val Val Val Ser Ile Asn Pro Gly Ala Gly Ser Gln Val
 245 250 255

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Asp Ala Gly Ser Thr Val Thr Ile Thr Thr Lys Ala Gly Ser Thr Gly
 260 265 270

Gly Gly Thr Gly Thr Gly
 275

<210> 10

<211> 268

<212> PRT

<213> *Streptomyces coelicolor*

<400> 10

Ser Gly Gln Phe Thr Lys Val Pro Pro Leu Leu Ser Lys Thr Glu Ala
 1 5 10 15

Gln Ala Arg Asp Arg Leu Asp Asp Ala Gly Leu Asp Val Gly Lys Val
 20 25 30

Arg His Ala Tyr Ser Asp Thr Val Glu Arg Gly Lys Val Ile Ser Thr
 35 40 45

Asp Pro Gly Val Gly Asp Arg Ile Arg Lys Asn Asp Ser Val Ser Leu
 50 55 60

Thr Val Ser Asp Gly Pro Asp Thr Val Lys Leu Pro Asp Val Thr Gly
 65 70 75 80

Tyr Lys Leu Asp Lys Ala Arg Thr Leu Leu Glu Asp Glu Gly Leu Glu
 85 90 95

Pro Gly Met Val Thr Arg Ala Phe Ser Asp Glu Val Ala Arg Gly Phe
 100 105 110

Val Ile Ser Thr Lys Pro Gly Ser Gly Thr Thr Val Arg Ala Gly Ser
 115 120 125

Ala Val Ala Leu Val Val Ser Lys Gly Ser Pro Val Asp Val Pro Asp
 130 135 140

Val Thr Gly Asp Asp Leu Asp Glu Ala Arg Ala Glu Leu Glu Gly Ala
 145 150 155 160

Gly Leu Lys Val Lys Thr Ala Asp Glu Arg Val Asn Ser Glu Tyr Asp
 165 170 175

Ser Gly Arg Val Ala Arg Gln Thr Pro Glu Pro Gly Gly Arg Ala Ala
 180 185 190

Glu Gly Asp Thr Val Thr Leu Thr Val Ser Lys Gly Pro Arg Met Ile
 195 200 205

Glu Val Pro Asp Val Val Gly Asp Ser Val Asp Asp Ala Lys Gln Lys
 210 215 220

Leu Glu Asp Ala Gly Phe Glu Val Asp Glu Asp Arg Gly Leu Leu Gly
 225 230 235 240

Leu Phe Gly Asp Thr Val Lys Lys Gln Ser Val Asp Gly Gly Asp Thr

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245

250

255

Ala Pro Glu Gly Ser Thr Val Thr Ile Thr Ile Arg
260 265

<210> 11

<211> 279

<212> PRT

<213> Streptomyces coelicolor

<400> 11

Gly Asn Asp Lys Val Pro Val Pro Ala Phe Ile Gly Leu Ser Lys Ala
1 5 10 15

Asp Ala Gln Gln Gln Ala Asp Asn Ile Asp Leu Val Leu Thr Phe Lys
20 25 30

Gln Gln Glu Cys Glu Asp Gln Pro Lys Gly Asn Ile Cys Ala Gln Asp
35 40 45

Pro Lys Gln Gly Thr Asp Val Asp Lys Glu Ser Thr Val Asn Leu Val
50 55 60

Val Ser Thr Gly Ala Pro Lys Val Ala Val Pro Asn Val Ile Asp Lys
65 70 75 80

Asn Ile Asp Glu Ala Lys Lys Gln Leu Glu Asp Lys Gly Phe Glu Val
85 90 95

Glu Thr Lys Gln Thr Glu Ser Ser Gln Asp Glu Gly Thr Ile Leu Ser
100 105 110

Gln Asn Pro Asp Pro Gly Lys Glu Leu Glu Lys Gly Ser Thr Val Thr
115 120 125

Leu Glu Val Ala Lys Ala Glu Glu Lys Ala Thr Val Pro Asp Val Val
130 135 140

Gly Arg Thr Cys Asp Glu Ala Lys Ala Gln Val Glu Ser Gly Gly Asp
145 150 155 160

Leu Thr Ala Val Cys Thr Asp Gln Pro Thr Asn Asp Pro Asn Gln Val
165 170 175

Gly Lys Val Ile Ser Thr Thr Pro Gln Ser Ser Thr Gln Val Asp Pro
180 185 190

Gly Ser Lys Val Thr Ile Val Val Gly Lys Ala Val Glu Lys Thr Lys
195 200 205

Val Pro Glu Val Arg Gly Lys Thr Leu Ala Glu Ala Arg Gln Ile Leu
210 215 220

Gln Gln Ser Gly Phe Thr Asn Val Gln Val Ala Gln Gly Ser Pro Gly
225 230 235 240

Asp Asp Asn Ala Lys Val Phe Ala Ser Asn Pro Gln Pro Gly Ser Glu
245 250 255

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Val Asp Asp Pro Ala Ala Thr Pro Ile Thr Leu Met Thr Val Pro Gly
 260 265 270

Asp Gly Gly Asn Gly Asn Gly
 275

<210> 12

<211> 277

<212> PRT

<213> Bacillus subtilis

<400> 12

Met Pro Lys Asp Val Lys Ile Pro Asp Val Ser Gly Met Glu Tyr Glu
 1 5 10 15

Lys Ala Ala Gly Leu Leu Glu Lys Glu Gly Leu Gln Val Asp Ser Glu
 20 25 30

Val Leu Glu Ile Ser Asp Glu Lys Ile Glu Glu Gly Leu Met Val Lys
 35 40 45

Thr Asp Pro Lys Ala Asp Thr Thr Val Lys Glu Gly Ala Thr Val Thr
 50 55 60

Leu Tyr Lys Ser Thr Gly Lys Ala Lys Thr Glu Ile Gly Asp Val Thr
 65 70 75 80

Gly Gln Thr Val Asp Gln Ala Lys Lys Ala Leu Lys Asp Gln Gly Phe
 85 90 95

Asn His Val Thr Val Asn Glu Val Asn Asp Glu Lys Asn Ala Gly Thr
 100 105 110

Val Ile Asp Gln Asn Pro Ser Ala Gly Thr Glu Leu Val Pro Ser Glu
 115 120 125

Asp Gln Val Lys Leu Thr Val Ser Ile Gly Pro Glu Asp Ile Thr Leu
 130 135 140

Arg Asp Leu Lys Thr Tyr Ser Lys Glu Ala Ala Ser Gly Tyr Leu Glu
 145 150 155 160

Asp Asn Gly Leu Lys Leu Val Glu Lys Glu Ala Tyr Ser Asp Asp Val
 165 170 175

Pro Glu Gly Gln Val Val Lys Gln Lys Pro Ala Ala Gly Thr Ala Val
 180 185 190

Lys Pro Gly Asn Glu Val Glu Val Thr Phe Ser Leu Gly Pro Glu Lys
 195 200 205

Lys Pro Ala Lys Thr Val Lys Glu Lys Val Lys Ile Pro Tyr Glu Pro
 210 215 220

Glu Asn Glu Gly Asp Glu Leu Gln Val Gln Ile Ala Val Asp Asp Ala
 225 230 235 240

Asp His Ser Ile Ser Asp Thr Tyr Glu Glu Phe Lys Ile Lys Glu Pro
 245 250 255

Thr Glu Arg Thr Ile Glu Leu Lys Ile Glu Pro Gly Gln Lys Gly Tyr
260 265 270

Tyr Gln Val Met Val
275

<210> 13
<211> 16
<212> PRT
<213> Mycobacterium tuberculosis

<400> 13

Gly Ser Ser His His His His His His Ser Ser Gly Leu Val Pro Arg
1 5 10 15

<210> 14
<211> 28
<212> PRT
<213> Mycobacterium tuberculosis

<400> 14

Ala Ile Ala Asp Ser Gly Asn Ser Val Pro Gln Thr Ala Ala Val Ile
1 5 10 15

Gly Thr Ala Gln Tyr Leu Ser Pro Glu Gln Ala Arg
20 25

<210> 15
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> synthetic oligonucleotide

<400> 15
gatagccata tgaccacccc ttcc

24

<210> 16
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> synthetic oligonucleotide

<400> 16
aaaccgaagc ttaacggccc accg

24

<210> 17
<211> 22
<212> DNA
<213> artificial sequence

<220>

<223> synthetic oligonucleotide

<400> 17

cgggggcata tggcgcgcggt ga

22

<210> 18

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 18

gcagtcgtaa gcttatgccg ccg

23

<210> 19

<211> 469

<212> PRT

<213> Mycobacterium tuberculosis

<400> 19

Met Thr Thr Arg Leu Gln Ala Pro Val Ala Val Thr Pro Pro Leu Pro
 1 5 10 15

Thr Arg Arg Asn Ala Glu Leu Leu Leu Leu Cys Phe Ala Ala Val Ile
 20 25 30

Thr Phe Ala Ala Leu Leu Val Val Gln Ala Asn Gln Asp Gln Gly Val
 35 40 45

Pro Trp Asp Leu Thr Ser Tyr Gly Leu Ala Phe Leu Thr Leu Phe Gly
 50 55 60

Ser Ala His Leu Ala Ile Arg Arg Phe Ala Pro Tyr Thr Asp Pro Leu
 65 70 75 80

Leu Leu Pro Val Val Ala Leu Leu Asn Gly Leu Gly Leu Val Met Ile
 85 90 95

His Arg Leu Asp Leu Val Asp Asn Glu Ile Gly Glu His Arg His Pro
 100 105 110

Ser Ala Asn Gln Gln Met Leu Trp Thr Leu Val Gly Val Ala Ala Phe
 115 120 125

Ala Leu Val Val Thr Phe Leu Lys Asp His Arg Gln Leu Ala Arg Tyr
 130 135 140

Gly Tyr Ile Cys Gly Leu Ala Gly Leu Val Phe Leu Ala Val Pro Ala
 145 150 155 160

Leu Leu Pro Ala Ala Leu Ser Glu Gln Asn Gly Ala Lys Ile Trp Ile
 165 170 175

Arg Leu Pro Gly Phe Ser Ile Gln Pro Ala Glu Phe Ser Lys Ile Leu

180	185	190
Leu Leu Ile Phe Phe Ser Ala Val Leu Val Ala Lys Arg Gly Leu Phe 195 200 205		
Thr Ser Ala Gly Lys His Leu Leu Gly Met Thr Leu Pro Arg Pro Arg 210 215 220		
Asp Leu Ala Pro Leu Leu Ala Ala Trp Val Ile Ser Val Gly Val Met 225 230 235 240		
Val Phe Glu Lys Asp Leu Gly Ala Ser Leu Leu Leu Tyr Thr Ser Phe 245 250 255		
Leu Val Val Val Tyr Leu Ala Thr Gln Arg Phe Ser Trp Val Val Ile 260 265 270		
Gly Leu Thr Leu Phe Ala Ala Gly Thr Leu Val Ala Tyr Phe Ile Phe 275 280 285		
Glu His Val Arg Leu Arg Val Gln Thr Trp Leu Asp Pro Phe Ala Asp 290 295 300		
Pro Asp Gly Thr Gly Tyr Gln Ile Val Gln Ser Leu Phe Ser Phe Ala 305 310 315 320		
Thr Gly Gly Ile Phe Gly Thr Gly Leu Gly Asn Gly Gln Pro Asp Thr 325 330 335		
Val Pro Ala Ala Ser Thr Asp Phe Ile Ile Ala Ala Phe Gly Glu Glu 340 345 350		
Leu Gly Leu Val Gly Leu Thr Ala Ile Leu Met Leu Tyr Thr Ile Val 355 360 365		
Ile Ile Arg Gly Leu Arg Thr Ala Ile Ala Thr Arg Asp Ser Phe Gly 370 375 380		
Lys Leu Leu Ala Ala Gly Leu Ser Ser Thr Leu Ala Ile Gln Leu Phe 385 390 395 400		
Ile Val Val Gly Gly Val Thr Arg Leu Ile Pro Leu Thr Gly Leu Thr 405 410 415		
Thr Pro Trp Met Ser Tyr Gly Gly Ser Ser Leu Leu Ala Asn Tyr Ile 420 425 430		
Leu Leu Ala Ile Leu Ala Arg Ile Ser His Gly Ala Arg Arg Pro Leu 435 440 445		
Arg Thr Arg Pro Arg Asn Lys Ser Pro Ile Thr Ala Ala Gly Thr Glu 450 455 460		
Val Ile Glu Arg Val 465		

<210> 20

<211> 491

<212> PRT

<213> Mycobacterium tuberculosis

<400> 20

Met Asn Ala Ser Leu Arg Arg Ile Ser Val Thr Val Met Ala Leu Ile
 1 5 10 15
 Val Leu Leu Leu Leu Asn Ala Thr Met Thr Gln Val Phe Thr Ala Asp
 20 25 30
 Gly Leu Arg Ala Asp Pro Arg Asn Gln Arg Val Leu Leu Asp Glu Tyr
 35 40 45
 Ser Arg Gln Arg Gly Gln Ile Thr Ala Gly Gly Gln Leu Leu Ala Tyr
 50 55 60
 Ser Val Ala Thr Asp Gly Arg Phe Arg Phe Leu Arg Val Tyr Pro Asn
 65 70 75 80
 Pro Glu Val Tyr Ala Pro Val Thr Gly Phe Tyr Ser Leu Arg Tyr Ser
 85 90 95
 Ser Thr Ala Leu Glu Arg Ala Glu Asp Pro Ile Leu Asn Gly Ser Asp
 100 105 110
 Arg Arg Leu Phe Gly Arg Arg Leu Ala Asp Phe Phe Thr Gly Arg Asp
 115 120 125
 Pro Arg Gly Gly Asn Val Asp Thr Thr Ile Asn Pro Arg Ile Gln Gln
 130 135 140
 Ala Gly Trp Asp Ala Met Gln Gln Gly Cys Tyr Gly Pro Cys Lys Gly
 145 150 155 160
 Ala Val Val Ala Leu Glu Pro Ser Thr Gly Lys Ile Leu Ala Leu Val
 165 170 175
 Ser Ser Pro Ser Tyr Asp Pro Asn Leu Leu Ala Ser His Asn Pro Glu
 180 185 190
 Val Gln Ala Gln Ala Trp Gln Arg Leu Gly Asp Asn Pro Ala Ser Pro
 195 200 205
 Leu Thr Asn Arg Ala Ile Ser Glu Thr Tyr Pro Pro Gly Ser Thr Phe
 210 215 220
 Lys Val Ile Thr Thr Ala Ala Ala Leu Ala Ala Gly Ala Thr Glu Thr
 225 230 235 240
 Glu Gln Leu Thr Ala Ala Pro Thr Ile Pro Leu Pro Gly Ser Thr Ala
 245 250 255
 Gln Leu Glu Asn Tyr Gly Gly Ala Pro Cys Gly Asp Glu Pro Thr Val
 260 265 270
 Ser Leu Arg Glu Ala Phe Val Lys Ser Cys Asn Thr Ala Phe Val Gln
 275 280 285
 Leu Gly Ile Arg Thr Gly Ala Asp Ala Leu Arg Ser Met Ala Arg Ala
 290 295 300
 Phe Gly Leu Asp Ser Pro Pro Arg Pro Thr Pro Leu Gln Val Ala Glu

305		310		315		320
Ser Thr Val Gly	Pro Ile Pro Asp Ser	Ala Ala Leu Gly	Met Thr Ser			
	325	330	335			
Ile Gly Gln Lys	Asp Val Ala Leu Thr	Pro Leu Ala Asn	Ala Glu Ile			
	340	345	350			
Ala Ala Thr Ile	Ala Asn Gly Gly	Ile Thr Met Arg	Pro Tyr Leu Val			
	355	360	365			
Gly Ser Leu Lys	Gly Pro Asp Leu	Ala Asn Ile Ser	Thr Thr Val Gly			
	370	375	380			
Tyr Gln Gln Arg	Arg Ala Val Ser	Pro Gln Val Ala	Ala Lys Leu Thr			
	385	390	395			400
Glu Leu Met Val	Gly Ala Glu Lys	Val Ala Gln Gln	Lys Gly Ala Ile			
	405	410	415			
Pro Gly Val Gln	Ile Ala Ser Lys	Thr Gly Thr Ala	Glu His Gly Thr			
	420	425	430			
Asp Pro Arg His	Thr Pro Pro His	Ala Trp Tyr Ile	Ala Phe Ala Pro			
	435	440	445			
Ala Gln Ala Pro	Lys Val Ala Val	Ala Val Leu Val	Glu Asn Gly Ala			
	450	455	460			
Asp Arg Leu Ser	Ala Thr Gly Gly	Ala Leu Ala Ala	Pro Ile Gly Arg			
	465	470	475			480
Ala Val Ile Glu	Ala Ala Leu Gln	Gly Glu Pro				
	485	490				